

GenCore version 5.1.6
Copyright (c) 1993 - 2003 Compugen Ltd.

OM protein - protein search, using sw model

Run on: May 29, 2003, 15:17:03 ; Search time 27 Seconds
(without alignments)
88.269 Million cell updates/sec

Title: US-09-924-102-2

Perfect score: 418

Sequence: 1 MLSTHFLYFLYFLYFLYSL.....RMGGQGRGTADTGMEFLS 81

Scoring table:

BLOSUM62

Gapop 10.0 , Gapext 0.5

Searched: 262574 seqs, 29422922 residues

Total number of hits satisfying chosen parameters: 262574

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 45 summaries

Database :

Issued_Patents_AA:*
1: /cgn2_6/ptodata/1/1aa/5A.COMB.pep:*
2: /cgn2_6/ptodata/1/1aa/5B.COMB.pep:*
3: /cgn2_6/ptodata/1/1aa/6A.COMB.pep:*
4: /cgn2_6/ptodata/1/1aa/6B.COMB.pep:*
5: /cgn2_6/ptodata/1/1aa/PTCUTS.COMB.pep:*
6: /cgn2_6/ptodata/1/1aa/Backfile1.pep:*

Pred. No. is the number of results predicted by chance to have a
score greater than or equal to the score of the result being printed,
and is derived by analysis of the total score distribution.

SUMMARIES

Result	Query	Match	Length	DB	ID	Description
No.	Score					
1	65	15.6	65	1	US-08-123-343A-2	Sequence 2, Appli
2	65	15.6	222	4	US-09-605-785-479	Sequence 479, App
3	65	15.6	222	4	US-09-439-313-479	Sequence 479, App
4	64	15.3	175	4	US-09-395-689-4	Sequence 4, Appli
5	64	15.3	633	1	US-08-458-477A-5	Sequence 5, Appli
6	64	15.3	633	2	US-09-033-153-5	Sequence 5, Appli
7	64	15.3	633	2	US-09-325-450B-5	Sequence 5, Appli
8	64	15.3	765	2	US-08-663-112-2	Sequence 2, Appli
9	62	14.8	63	4	US-08-828-683A-28	Sequence 28, Appli
10	62	14.8	1898	1	US-08-056-200-94	Sequence 28, Appli
11	62	14.8	1898	2	US-08-800-644-94	Sequence 94, Appli
12	61.5	14.7	806	1	US-07-980-528-2	Sequence 2, Appli
13	60	14.4	127	4	US-08-525-539A-65	Sequence 65, Appli
14	59.5	14.2	151	4	US-08-858-207A-475	Sequence 475, App
15	59.5	14.2	399	4	US-09-222-938A-49	Sequence 49, Appli
16	59.5	14.2	595	4	US-09-370-838-187	Sequence 187, App
17	59.5	14.2	760	1	US-08-195-152-2	Sequence 2, Appli
18	58.5	14.0	2273	4	US-09-426-998-5	Sequence 2, Appli
19	58	13.9	1958	1	US-07-945-283-2	Sequence 28, Appli
20	57.5	13.8	126	2	US-08-822-028-28	Sequence 28, Appli
21	57.5	13.8	126	2	US-08-479-285-28	Sequence 28, Appli
22	57	13.6	29	1	US-07-694-983-14	Sequence 14, Appli
23	57	13.6	3169	2	US-08-477-451-6	Sequence 6, Appli
24	56	13.4	105	1	US-08-276-852-93	Sequence 93, Appli
25	56	13.4	105	1	US-08-899-575-93	Sequence 93, Appli
26	56	13.4	105	1	US-08-899-575-93	Sequence 93, Appli
27	56	13.4	105	5	PCT-US95-08743-93	Sequence 93, Appli

28	55	13.2	129	4	US-08-943-136-2	Sequence 2, Appli
29	55	13.2	129	3	US-08-973-518-2	Sequence 2, Appli
30	54.5	13.0	239	4	US-08-812-586-29	Sequence 29, Appli
31	54.5	13.0	449	1	US-07-917-722-2	Sequence 2, Appli
32	54.5	13.0	449	2	US-08-489-666C-3	Sequence 3, Appli
33	54.5	13.0	449	2	US-08-911-092-3	Sequence 3, Appli
34	54.5	13.0	449	2	US-08-485-001B-3	Sequence 3, Appli
35	54.5	13.0	449	3	US-08-454-121A-3	Sequence 3, Appli
36	54.5	13.0	449	4	US-08-482-161B-3	Sequence 3, Appli
37	54.5	13.0	449	4	US-09-057-963A-2	Sequence 2, Appli
38	54.5	13.0	577	4	US-09-261-855-2	Sequence 2, Appli
39	54	12.9	266	1	US-08-247-809A-12	Sequence 12, Appli
40	54	12.9	266	1	US-08-711-728-12	Sequence 12, Appli
41	54	12.9	318	1	US-08-247-809A-2	Sequence 2, Appli
42	54	12.9	318	1	US-08-247-809A-6	Sequence 6, Appli
43	54	12.9	318	2	US-08-711-728-2	Sequence 2, Appli
44	54	12.9	318	2	US-08-711-728-6	Sequence 6, Appli
45	54	12.9	410	1	US-08-153-848-7	Sequence 7, Appli

ALIGNMENTS

RESULT 1
US-08-123-343A-2
Sequence 2, Application US/08123343A
Patent No. 5593879
GENERAL INFORMATION:
APPLICANT: Steller, Hermann
APPLICANT: Abrams, John M.
APPLICANT: Grether, Megan E.
APPLICANT: White, Kristin
TITLE OF INVENTION: Cell Death Genes of Drosophila
TITLE OF INVENTION: Melanogaster and Vertebrate Analogs
NUMBER OF SEQUENCES: 16
CORRESPONDENCE ADDRESS:
ADDRESS: Hamilton, Brook, Smith & Reynolds, P.C.
STREET: Two Millitia Drive
CITY: Lexington
STATE: MA
COUNTRY: US
ZIP: 02173
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/123, 343A
FILING DATE: 17-SEP-1993
CLASSIFICATION: 800
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/004,957
FILING DATE: 15-JAN-1993
ATTORNEY/AGENT INFORMATION:
NAME: Granahan, Patricia
REGISTRATION NUMBER: 32,227
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-861-6240
TELEFAX: 61861-9540
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
LENGTH: 65 amino acids
TYPE: amino acid
STRANDEDNESS: unknown
TOPOLOGY: unknown
MOLECULE TYPE: protein
US-08-123-343A-2
Query Match 15.6%; Score 65; DB 1; Length 65;
Best Local Similarity 37.7%; Pred. No. 0.36;
Matches 23; Conservative 11; Mismatches 11; Indels 16; Gaps 4;

QY 16 LSYSLGDRARLCLRTKQOQEQOILROSEVLEF-----SETLR-----KTGKGR 62
Db 3 VAFYIPQATL-LREA--EQEQOILRLRESQWFLATVYLELRLQTSCHPRTGRSGK 59
QY 63 W 63
Db 60 Y 60

RESULT 2

US-09-605-785-479
; Sequence 479, Application US/09605785
; Patent No. 6321716
; GENERAL INFORMATION:
; APPLICANT: Xu, Jiangchun
; APPLICANT: Dillon, Davin C.
; APPLICANT: Mitcham, Jennifer L.
; APPLICANT: Harlocker, Susan L.
; APPLICANT: Jiang, Yuqi
; APPLICANT: Henderson, Robert A.
; APPLICANT: Kalos, Michael D.
; APPLICANT: Fanger, Gary R.
; APPLICANT: Retter, Marc W.
; APPLICANT: Stolk, John A.
; APPLICANT: Day, Craig H.
; APPLICANT: Vedvick, Thomas S.
; APPLICANT: Carter, Darrick
; APPLICANT: Li, Samuel
; APPLICANT: Wang, Aijun
; APPLICANT: Skeiky, Yasir A.W.
; APPLICANT: Hepler, William
; TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND
; FILE REFERENCE: 210121.427C16
; CURRENT APPLICATION NUMBER: US/09/605,785
; CURRENT FILING DATE: 2000-06-27
; NUMBER OF SEQ ID NOS: 835
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 479
; LENGTH: 222
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-605-785-479

Query Match 15.6%; Score 65; DB 4; Length 222;
Best Local Similarity 52.0%; Pred. No. 1.7;
Matches 13; Conservative 6; Mismatches 2; Indels 4; Gaps 1;

QY 11 LFTYPLSYSGDRARLCLRTKQOQ 35
Db 199 IFYF---LGNQARLCLKRRKKQ 219

RESULT 3
US-09-439-313-479
; Sequence 479, Application US/09439313
; Patent No. 6329505
; GENERAL INFORMATION:
; APPLICANT: Xu, Jiangchun
; APPLICANT: Dillon, Davin C.
; APPLICANT: Mitcham, Jennifer L.
; APPLICANT: Harlocker, Susan Louise
; APPLICANT: Jiang, Yuqi
; APPLICANT: Reed, Steven G.
; APPLICANT: Kalos, Michael
; APPLICANT: Fanger, Gary
; APPLICANT: Retter, Marc
; APPLICANT: Stolk, John
; APPLICANT: Day, Craig
; TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THERAPY AND
; FILE REFERENCE: 210121.427C9

; CURRENT APPLICATION NUMBER: US/09/439,313
; CURRENT FILING DATE: 1999-11-12
; NUMBER OF SEQ ID NOS: 575
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 479
; LENGTH: 222
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-439-313-479

Query Match 15.6%; Score 65; DB 4; Length 222;
Best Local Similarity 52.0%; Pred. No. 1.7;
Matches 13; Conservative 6; Mismatches 2; Indels 4; Gaps 1;

QY 11 LFTYPLSYSGDRARLCLRTKQOQ 35
Db 199 IFYF---LGNQARLCLKRRKKQ 219

RESULT 4
US-09-395-689-4
; Sequence 4, Application US/09395689
; Patent No. 6387684
; GENERAL INFORMATION:
; APPLICANT: Hwang, Jaulang
; APPLICANT: Hui, Cho-Fat
; APPLICANT: Chen, Tzong-Yueh
; TITLE OF INVENTION: TOPOISOMERASE 1-MEDIATED DNA DELIVERY
; FILE REFERENCE: 089191/024001
; CURRENT APPLICATION NUMBER: US/09/395,689
; CURRENT FILING DATE: 1999-09-13
; NUMBER OF SEQ ID NOS: 6
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 4
; LENGTH: 175
; TYPE: PRT
; ORGANISM: Pseudomonas aeruginosa
US-09-395-689-4

Query Match 15.3%; Score 64; DB 4; Length 175;
Best Local Similarity 30.0%; Pred. No. 1.7;
Matches 18; Conservative 9; Mismatches 33; Indels 0; Gaps 0;

QY 18 YSLGDRARLCLRTKQOQEQOILROSEVLEFSETLRKTGKGRWGCGGGRGTADTG 77
Db 23 YETARRLKCVDRIRNQYREDWKSKEKVKQRAVALYFTDKLALRAGNEKEEGTADTVG 82

RESULT 5
US-08-458-477A-5
; Sequence 5, Application US/08458477A
; Patent No. 5723311
; GENERAL INFORMATION:
; APPLICANT: WEI, ET AL.
; TITLE OF INVENTION: Human Topoisomerase I
; NUMBER OF SEQUENCES: 5
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: CARELLA, BYRNE, BAIN, GUILFILLAN,
; ADDRESSEE: CECCHI, STEWART & OLSTEIN
; STREET: 6 BECKER FARM ROAD
; CITY: ROSELAND
; STATE: NEW JERSEY
; COUNTRY: USA
; ZIP: 07068
; COMPUTER READABLE FORM:
; MEDIUM TYPE: 3.5 INCH DISKETTE
; COMPUTER: IBM PS/2
; OPERATING SYSTEM: MS-DOS
; SOFTWARE: WORD PERFECT 5.1
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/458,477A
; FILING DATE: June 2, 1995
; CLASSIFICATION: 435

PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/US94/05701
FILING DATE: 18 MAY 94
ATTORNEY/AGENT INFORMATION:
NAME: MULLINS, J.G.
REGISTRATION NUMBER: 33,073
REFERENCE/DOCKET NUMBER: 325800-309 (PFI18P1)
TELECOMMUNICATION INFORMATION:
TELEPHONE: 201-994-1700
TELEFAX: 201-994-1744
INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
LENGTH: 633 AMINO ACIDS
TYPE: AMINO ACID
STRANDEDNESS:
TOPOLOGY: LINEAR
MOLECULE TYPE: PROTEIN
US-08-458-477A-5

Query Match: 15.3%; Score 64; DB 1; Length 633;
Best Local Similarity 30.0%; Pred. No. 7.8;
Matches 18; Conservative 9; Mismatches 33; Indels 0; Gaps 0;

Qy 18 YSLGDRARLCLRTKQOQKEQOILROSEVLFERSETLRTKTKGKGRMGOGGGRGTADTGG 77
Db 312 YETARRLKVCVDKIRNQRREDWKSKEKVKRQRAVALYFDIKLALRAGNEKEGETADTVG 371

RESULT 6
US-09-033-153-5
Sequence 5, Application US/09033153
Patent No. 5968803

GENERAL INFORMATION:

APPLICANT: WEI, ET AL.
TITLE OF INVENTION: Human Topoisomerase I

NUMBER OF SEQUENCES: 5

CORRESPONDENCE ADDRESSES:

ADDRESSEE: CARELLA, BYRNE, BAIN, GILFILLAN,
STREET: 6 BECKER FARM ROAD
CITY: ROSELAND
STATE: NEW JERSEY
COUNTRY: USA
ZIP: 07068

COMPUTER READABLE FORM:

MEDIUM TYPE: 3.5 INCH DISKETTE

COMPUTER: IBM PS/2

OPERATING SYSTEM: MS-DOS

SOFTWARE: WORD PERFECT 5.1

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/09/033,153

FILING DATE:

CLASSIFICATION: 435

PRIOR APPLICATION DATA:

APPLICATION NUMBER: 08/458,477

FILING DATE: June 2, 1995

APPLICATION NUMBER: PCT/US94/05701

FILING DATE: 18 MAY 94

ATTORNEY/AGENT INFORMATION:

NAME: MULLINS, J.G.

REGISTRATION NUMBER: 33,073

REFERENCE/DOCKET NUMBER: 325800-309 (PFI18P1)

TELECOMMUNICATION INFORMATION:

TELEPHONE: 201-994-1700

TELEFAX: 201-994-1744

INFORMATION FOR SEQ ID NO: 5:

SEQUENCE CHARACTERISTICS:

LENGTH: 633 AMINO ACIDS

TYPE: AMINO ACID

STRANDEDNESS:

TOPOLOGY: LINEAR

MOLECULE TYPE: PROTEIN

US-09-033-153-5

Query Match: 15.3%; Score 64; DB 2; Length 633;
Best Local Similarity 30.0%; Pred. No. 7.8;
Matches 18; Conservative 9; Mismatches 33; Indels 0; Gaps 0;

Qy 18 YSLGDRARLCLRTKQOQKEQOILROSEVLFERSETLRTKTKGKGRMGOGGGRGTADTGG 77
Db 312 YETARRLKVCVDKIRNQRREDWKSKEKVKRQRAVALYFDIKLALRAGNEKEGETADTVG 371

RESULT 7
US-09-325-430B-5
Sequence 5, Application US/09325430B
Patent No. 6255077

GENERAL INFORMATION:

APPLICANT: Wei et al.

TITLE OF INVENTION: Human DNA Topoisomerase 1 Alpha

FILE REFERENCE: PFI18D2

CURRENT APPLICATION NUMBER: US/09/325,430B

CURRENT FILING DATE: 1999-06-04

PRIOR APPLICATION NUMBER: 09/033,153

PRIOR FILING DATE: 1998-03-02

PRIOR APPLICATION NUMBER: 08/458,477

PRIOR FILING DATE: 1995-06-02

PRIOR APPLICATION NUMBER: PCT/US94/05701

PRIOR FILING DATE: 1994-05-18

NUMBER OF SEQ ID NOS: 5

SOFTWARE: PatentIn Ver. 2.1

SEQ ID NO 5

LENGTH: 633

TYPE: PRT

ORGANISM: Homo sapiens

US-09-325-430B-5

Query Match: 15.3%; Score 64; DB 4; Length 633;
Best Local Similarity 30.0%; Pred. No. 7.8;
Matches 18; Conservative 9; Mismatches 33; Indels 0; Gaps 0;

Qy 18 YSLGDRARLCLRTKQOQKEQOILROSEVLFERSETLRTKTKGKGRMGOGGGRGTADTGG 77
Db 312 YETARRLKVCVDKIRNQRREDWKSKEKVKRQRAVALYFDIKLALRAGNEKEGETADTVG 371

RESULT 8

US-08-663-112-2

Sequence 2, Application US/08663112

Patent No. 5849503

GENERAL INFORMATION:

APPLICANT: WAGATSUMA, Masako

APPLICANT: KURITA, No. 58495031ko

TITLE OF INVENTION: MUTANT PROTEINS OF HUMAN DNA

TITLE OF INVENTION: TOPOISOMERASE I

NUMBER OF SEQUENCES: 7

CORRESPONDENCE ADDRESSES:

ADDRESSEE: Finnegan, Henderson, Farabow, Garrett &

ADDRESSEE: Dunner L.L.P.

STREET: 1300 I Street, N.W.

CITY: Washington

STATE: D.C.

COUNTRY: USA

ZIP: 20005-3315

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: PatentIn Release #1.0, Version #1.30

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/663,112

FILING DATE: 26-NOV-1996

CLASSIFICATION: 435

ATTORNEY/AGENT INFORMATION:

NAME: Elnaudi, Carolyn P.

REGISTRATION NUMBER: 32,220

STREET: 620 Newport Center Drive, Sixteenth Floor
CITY: Newport Beach
STATE: CA
COUNTRY: U.S.A.
ZIP: 92660
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/800,644
FILING DATE: 14-FEB-1997
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/056,200
FILING DATE: 30-APR-1993
ATTORNEY/AGENT INFORMATION:
NAME: Fedrick, Michael F.
REGISTRATION NUMBER: 36,799
REFERENCE/DOCKET NUMBER: NIH054,001A
TELECOMMUNICATION INFORMATION:
TELEPHONE: (714) 760-0404
TELEFAX: (714) 760-9502
INFORMATION FOR SEQ ID NO: 94:
SEQUENCE CHARACTERISTICS:
LENGTH: 1898 amino acids
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
US-08-800-644-94

Query Match 14.8%; Score 62; DB 2; Length 1898;
Best Local Similarity 31.7%; Pred. No. 52;
Matches 13; Conservative 13; Mismatches 15; Indels 0; Gaps 0;
QY 22 DRARLCRTKQOQKLEQILROSEVLFPSRTLRKTKGKGR 62
DB 333 ERREQQLRRQEQERREQQLRRQEQERREQQLRRQEQER 373

RESULT 12
US-07-980-528-2
Sequence 2, Application US/07980528
Patent No. 5457026
GENERAL INFORMATION:
APPLICANT: Dreyfuss, Gideon
APPLICANT: Kiledjian, Megeditch
TITLE OF INVENTION: METHODS OF PROMOTING INTERMOLECULAR
TITLE OF INVENTION: INTERACTIONS INVOLVING A NUCLEIC ACID AND MOLECULES
TITLE OF INVENTION: USERFL
NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
ADDRESSEE: Woodcock Washburn Kurtz Mackiewicz &
ADDRESSEE: No. 5457026rls
STREET: One Liberty Place 46th Floor
CITY: Philadelphia
STATE: Pennsylvania
COUNTRY: USA
ZIP: 19403
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentn Release #1.0, Version #1.25; mb/MD
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/980,528
FILING DATE: 19921120
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Deluca, Mark

REGISTRATION NUMBER: 33,229
REFERENCE/DOCKET NUMBER: UPN-0847
TELECOMMUNICATION INFORMATION:
TELEPHONE: (215) 568-3100
TELEFAX: (215) 568-3439
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
LENGTH: 806 amino acids
TYPE: AMINO ACID
TOPOLOGY: linear
MOLECULE TYPE: protein
US-07-980-528-2

Query Match 14.7%; Score 61.5; DB 1; Length 806;
Best Local Similarity 30.2%; Pred. No. 21;
Matches 19; Conservative 7; Mismatches 26; Indels 11; Gaps 1;

QY 28 LRKTKOQKLEQILROSEVLFPSRTLRKTKGK-----RRMGOGGGRGTADTG 76
DB 638 LQKEAQKLEQYEESEKALPPEKQNTGSKSNKNSGKNQPNRGGRHGRGLMMRG 697
QY 77 GMF 79
DB 698 GNF 700

RESULT 13
US-08-525-539A-65
Sequence 65, Application US/08525539A
Patent No. 6309636
GENERAL INFORMATION:
APPLICANT: DO COUTO, FERNANDO J.R.
APPLICANT: CERTANT, ROBERTO L.
APPLICANT: PETERSON, JERRY A.
TITLE OF INVENTION: RECOMBINANT PEPTIDES DERIVED FROM THE
TITLE OF INVENTION: M33 ANTI-BA46 ANTIBODY, METHODS OF USE THEREOF, AND
TITLE OF INVENTION: METHODS OF HUMANIZING ANTIBODY PEPTIDES
NUMBER OF SEQUENCES: 81
CORRESPONDENCE ADDRESS:
ADDRESSEE: MORRISON & FOERSTER
STREET: 755 Page Mill Road
CITY: Palo Alto
STATE: CA
COUNTRY: USA
ZIP: 94304-1018
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/525,539A
FILING DATE: 14-SEP-1995
CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: DYLAN, TYLER
REGISTRATION NUMBER: 37,612
REFERENCE/DOCKET NUMBER: 27633-20001.21
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 813-5600
TELEFAX: (415) 494-0792
TELEX: 706141
INFORMATION FOR SEQ ID NO: 65:
SEQUENCE CHARACTERISTICS:
LENGTH: 127 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
US-08-525-539A-65

Query Match 14.4%; Score 60; DB 4; Length 127;
Best Local Similarity 24.2%; Pred. No. 3.6;
Matches 22; Conservative 15; Mismatches 28; Indels 26; Gaps 3;

